

Table 1

SEQUENCE IDENTIFICATION NUMBERS

SEQ ID NO	NAME	NS	AS	Sequence
1	LT 1-1	60	20	WLGTFSGTCSTAFYFPLGVP
2	LT 6-1	60	20	CRWLRTKRTLPLFSVMPFWC
3	LT 7-1	61	31	MYSNVSDVAADGVSCCVCCSWVQNDRPDSG*
4	LT 8-2	60	20	YKWRMGVYLSGVRLMRAFI
5	LT 9-1	60	20	VSYRTCTAGGQMSRWRLFII
6	LT 12-1	60	20	GYSLTSMASAFVRPCVCGSL
7	LT 13-1	60	20	WLGTFSGTCSTAFYFPLGVP
8	LT 15-2	60	20	TNFRPSPTFHAILLWPNTFS
9	LT 16-1	60	20	VGLGGWCFCDCYVVAWDFQTQ
10	LT 19-2	60	20	FDYCGPLICLHCNLGRCVS
11	LT 21-2	61	5	MMDSA*
12	LT 22-2	60	20	ASRLYPIGGTVPFGRGTGAG
13	LT 23-2	60	20	GDYGCCWVVTGTGVCRCYVW
14	LT 27-1	60	20	ACWALWSLFRQDLLLLVITFD
15	LT 28-1	61	31	DSAPGERYFVDFLGVSFACVWSVQNDRPDSG*
16	LT 29-1	60	14	IPWAPPMYFADSNV*
17	LT 33-1	60	20	TPSCRAGVLRCTGCFGVRSG
18	LT 34-2	60	20	LWRCRTVSAYLSWLRHYSSS
19	LT 35-1	60	20	HSRPALCMVSLRWARSLWIV
20	LT 37-2	42	14+?	WTHVWVGWL VAGMS
21	LT 38-1	60	20	RFRCRADLCVTLTVLSFLAQ

Table 1, DK62021PC

WO 2004/003008

PCT/EP2003/006958

22	LT 45-2	60	20	CLETLRVCPYVARIAIQHLR
23	LT 48-3	60	20	LLAWRVQQSRPLPYLHIAFI
24	LT 49-2	60	20	PPPLTGRWSRQCVSFVGVH
25	LT 54-1	60	20	CWHRAWMLSWHGVWSLTIV
26	LT 62-1	60	20	APPIGRWRGLYMRSRFVSL
27	LT 76-3	60	18	VRLFIVCIHCCLMLLVG*
28	LT 79-1	60	20	IPSCSVLVCLCHLARLWHCE
29	LT 82-3	60	20	CSVMHVFRVGPSSGSLSCG
30	LT 93-1	126	42	RATYWFRSRYQVVHRSRLPYGPLIVVGIGALNLELNRTLCS
31	LT 103-1	60	20	SLAIWSTQSCARCQCTLSRV
32	LT 105-2	60	20	FWFLPAPPCKCGLLYRLSVH
33	LT 108-1	60	20	LAGRHFSRVVDRIRYRLWT
34	LT 126-2	60	20	MPIPLCRSAGRLLYLYTHY
35	LT 127-1	60	20	YTLPVLLCLMRTGMLRCAC
36	LT 135-2	60	20	GMPIRASPCYLGVDGWXCXL
37	LT 138-2	60	20	KPWEYLRMPFWMRVARFFIW
38	LT 140-2	60	20	ALLMFGCPNWFASWRHLFI
39	ND	ND	ND	EFGSGLVSGRGIIVRRMLFLRVLLWVLRSAEYES*
40	ND	ND	ND	GPFENWRVEELARGRYRMHGDVVLRS AEYES*
41	ND	ND	ND	GPIDCHILLWYSRQQRGGSGGP*
42	ND	ND	ND	GSRFRVFVCSLFSLSGRGGVVVLRSAEYES*
43	ND	ND	ND	GPFKRCHERLVAFARCWFMWSMVLRSAEYES*
44	ND	ND	ND	GPSNDNQLVLRVRLRVLIIVMRVLRSAEYES*
45	ND	ND	ND	EFVRRRMRLLVRLMGSDSDGTIPDFGP*
46	ND	ND	ND	GPSLQFLEVVSVCYMLVYDLSKGP*
47	ND	ND	ND	GPQPFCSPPSFYTRLLIIVRLLSLDLQRSSNRRY*
48	ND	ND	ND	GPAPLSLCVCKCGCGHTRPFVGP*
49	ND	ND	ND	GPDVHIWQSHIFYAMRHHMGP*

Table 1, DK62021PC

WO 2004/003008

PCT/EP2003/006958

50	ND	ND	ND	ND	ND	EFGSGCGFVRGRVRCVILLRVLRS AEIYES*
51	ND	ND	ND	ND	ND	GPHSSAHDRIWLVRGLRIULLVLRSAE IYES*
52	ND	ND	ND	ND	ND	EFGSGLCVRRWWGMSVGSRIMLVMLVLRSAE IYES*
53	ND	ND	ND	ND	ND	GPVYSEAFVCLVCAGVCVEECGGSLDLQRSSNR RY*
54	ND	ND	ND	ND	ND	GPIETVGFIVRLHTLLMVLRRRTGP*
55	ND	ND	ND	ND	ND	GPLHRTLVDMMCCWLMLESNMGP*
56	ND	ND	ND	ND	ND	EFGVRVVCVRSFLVRLCGLLRCRGVLRSAE IYES*
57	ND	ND	ND	ND	ND	EFVRECSLCRVMVLMFVLRGIRLRLVLRSAE IYES*
58	ND	ND	ND	ND	ND	EFGVRLLVLLRLRCVRRGGGCFVCWVLRSAE IYES*
59	ND	ND	ND	ND	ND	GSGFRMRVLVMVCLRVVFLVRRVLRSAE IYES*
60	ND	ND	ND	ND	ND	GRLGWRLLCVRIVLVCRLRGLVLRSAE IYES*
61	ND	ND	ND	ND	ND	EFGSGWYVDLGDYSVWVDYVYCGSGSLDLQRSSNR RY*
62	ND	ND	ND	ND	ND	GPSQCAQRVALIQMYIDALVCIGP*
63	ND	ND	ND	ND	ND	EFGSGCVIRVGVIRRRMLXLRFVFLVLRSAE IYES*
64	ND	ND	ND	ND	ND	GPLDPMLRQRFSLRVWIMLLGVLRSAE IYES*
65	ND	ND	ND	ND	ND	GPNSWVWRYVTIAHWLANYRMSGP*
66	ND	ND	ND	ND	ND	GPAMKSCTIRVFRVCIVLRVLRVLRSAE IYES*
67	ND	ND	ND	ND	ND	EFGVRMRMLRIRFRGLFVLRGFRGLVLRSAE IYES*
68	ND	ND	ND	ND	ND	GPVPSSPCSELLYCRDVLCHWPGP*
69	ND	ND	ND	ND	ND	GPCEPFIGDCWPCLIRTLVTLRGLDL
70	ND	ND	ND	ND	ND	GPWWKDRGVLVRLCVLRLVVGVVLRSAE IYES*
71	ND	ND	ND	ND	ND	GPRLVVRMRGWCRVSLJXFWLRVLRSAE IYES*
72	ND	ND	ND	ND	ND	EFRVRIUVSLRIWRLLVRRRCCLCLVLRSAE IYES*
73	ND	ND	ND	ND	ND	GPVECADVLEFASRIKLLCLCFRVLRS AEIYES*
74	ND	ND	ND	ND	ND	EFGRRLLVFRLSVFVVVLGRRLSRVLRSAE IYES*
75	ND	ND	ND	ND	ND	GAGLGRVIRLIVVLRICIFLLFRVLRSAE IYES*
76	ND	ND	ND	ND	ND	GPFPFDYPRWIMIVLLRGVLRSAE IYES*
77	ND	ND	ND	ND	ND	GSRGLRLCLLGRCLCGCLIMRVLRSAE IYES*

Table 1, DK62021PC

WO 2004/003008

PCT/EP2003/006958

78	ND	ND	ND	ND	GPESYVLPARGEALYYLRAWLGP*
79	ND	ND	ND	ND	GSRCIRRRISILFFVFRVLRSSRVLRSAEIYES*
80	ND	ND	ND	ND	GPFEHARGHVVTICRLRLFWLLRSAEIYES*
81	ND	ND	ND	ND	GPSSLLRRCLILGMVLGVLRRLRVLRSAEIYES*
82	ND	ND	ND	ND	GPHPVLA VQLINAYLGLERVGRGP*
83	ND	ND	ND	ND	GPLPSGAVSTEAYFWEVFKLLMGP*
84	ND	ND	ND	ND	GPYPYLRILLVQKIACVRRRALWVLRSAEIYES*
85	ND	ND	ND	ND	GPVGVEGVDSVFGWCVCFLVWSLDLQRSSNRRY*
86	ND	ND	ND	ND	EFRVRVLGCMGVFLRLRFCGGLRLRVLRSAEIYES*

Table 2:

Number	Peptide	Sequence
1	1	GSGLVSGRGIIVRRMLFLRVLLWVLRSAEIYES*
2	3	GPFENWRVEELARGRYRMHGDVVLRSAEIYES*
3	4	GPIDCIIFLLWYSRQQRGSGRGGP*
4	5	GSRFRVVFVCSLFSFLSGRGGGVVLRSAEIYES*
5	7	GPFKRCHERLVAFAFCWFMWSMVLRSAEIYES*
6	8	GPSNDNQLVLRVRILRVLIVMRVLRS AEIYES*
7	9	RVRRMRLLVRLMGSDDSGTIPDFGP*
8	10	GPSLQFLEVVS CYMVLYDLSKGP*
9	11	GPQPFCSPPSFYTRLLIIVRLLSLDLQRSSNRRY*
10	13	GPAPLSLCVCKCGCGHTRPFVGP*
11	14	GPDVHIWQSIIFYAMRHMMGP*
12	15	GSGCGCFVRGRIVRIRC VILLRVLRSAEIYES*
13	16	GPHSSAHDRIWLRVRGLRIILLVLRSAEIYES*
14	17	GSGLCVRRWWGMSVGSRI MLV MLVLRSAEIYES*
15	18	GPVYSEAFVCLVCAGVCVEECGGS L DLQRSSNRRY*
16	19	GPIETVGFI VRLHTLLMVLRTGP*
17	20	GPLHRTLLVDMCCWLMSLESNMGP*
18	22	GVRVVCVVRSLFVLRCGLLRGVLRS AEIYES*
19	23	VRECSLCRVMVLMFVLRGIRLVRSAEIYES*
20	28	GVRLLVLLRLRCVRRGGGCFVCWVLRSAEIYES*
21	30	GSGFRMRVLVMVCRLRVVFLVRRVLRSAEIYES*
22	33	GRLGWLRLLCVRIVLVCLRRGLVLRSAEIYES*
23	34	GSGWYVDLGDYSVWVDYVYCGSGS L DLQRSSNRRY*
24	37	GPSQCAQRVALIQMYIDALVCIGP*
25	40	GSGCVRIRVGIVRRMLFLRFVFLVP*
26	41	GPLLDPMRLRQRFSLRVWIMLLGVLRSAEIYES*
27	48	GPNSWVWRYVTIAHWLANYRMSGP*
28	49	GPAMKSCTIRVFRVCIVLRIVRVLRS AEIYES*
29	51	GVRMRLMIRIFRGLFVLRGFRGLVLRSAEIYES*
30	52	GPVPSSPCSFLLYCRDVLCHWPGP*
31	53	GPCEPFIGDCWPCLIRTLVTLRGLDLQRSSNRRY*
32	54	GPWWKDRGVLVRLCVLRLVVG VVLRSAEIYES*
33	57	GPRLLVRMRGWCRVSLXFWLRVLRSAEIYES*
34	59	RVRIIVVSLRIWRLLVRRRC LCLVLRSAEIYES*
35	60	GPVECADVLFASRI RLLCLCFRVLRS AEIYES*
36	62	GRRLLVFRLSVFVVVLGRRLSRVLRSAEIYES*
37	65	GAGLGRVIRLRIVVLR CIFLLFRVLRSAEIYES*
38	69	GPFPFDYPRWIMIVLLRGVLRSAEIYES*
39	72	GSRGLRLCCLGRCLCGCLIMRVLRSAEIYES*
40	74	GPESYVLWPARGEALYYLRAWLGP*
41	75	GSRCIRRRISILFFVFRVLR SRRVLRSAEIYES*
42	77	GPFSEHARGHVVTICRLRL LFVLLRS AEIYES*
43	79	GPSSLLRRCLILGMVLGVLR RRVLRSAEIYES*
44	88	GPHPVLA VQLINAYLGLERVGRGP*
45	89	GPLPSGAVSTEAYFWEVFKLLMGP*
46	90	GPYPYLRILLVQKIACVRRALWVLRSAEIYES*

Table 3:

(linear) peptide	Survivin	c-IAP-1	c-IAP-2	XIAP	Survivin	HPV16 E6
5	+	-	-	-	-	-
7	+	-	-	-	-	-
15	+	-	-	-	-	-
16	+	-	-	-	-	-
40	+	-	-	-	-	-
54	+	-	-	-	-	-
65	+	-	-	-	-	-
69	+	-	-	-	-	-
75	+	-	-	-	-	-
77	+	-	-	+	-	-
90	+	-	-	-	-	-
K1	-	-	-	-	-	-